

San-Hoffman

CRF Errors Corrected by the STIC Systems Branch

1642

P#11

Serial Number: 09/095,478A

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

RECEIVED

JUN 26 2000

TECH CENTER 1600/2900

*Examiner: ~~The above~~ corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/095,478ADATE: 06/08/2000
TIME: 07:09:25

INPUT SET: S35575.raw

<p>This Raw Listing contains the General Information Section and up to the first 5 pages.</p>

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SEQUENCE LISTING

(1) General Information:

(i) APPLICANT: Gregory Plowman
Bahija Jallal

(ii) TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF
PTP RELATED DISORDERS

(iii) NUMBER OF SEQUENCES: 23

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Brobeck, Phleger & Harrison LLP
(B) STREET: 12390 El Camino Real
(C) CITY: San Diego
(D) STATE: California
(E) COUNTRY: U.S.A.
(F) ZIP: 92130-2081

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
storage
(B) COMPUTER: IBM Compatible
(C) OPERATING SYSTEM: IBM P.C. DOS 5.0
(D) SOFTWARE: FastSEQ for Windows 2.0

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: 09/095,478
(B) FILING DATE: June 10, 1998
(C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: 60/049,756
(B) FILING DATE: June 11, 1997

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/095,478ADATE: 06/08/2000
TIME: 07:09:26

INPUT SET: S35575.raw

47

48

(A) APPLICATION NUMBER:

49

(B) FILING DATE:

50

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52

(viii) ATTORNEY/AGENT INFORMATION:

53

54

(A) NAME: Warburg, Richard J.

55

(B) REGISTRATION NUMBER: 32,327

56

(C) REFERENCE/DOCKET NUMBER: 235/054

57

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(ix) TELECOMMUNICATION INFORMATION:

63

64

(A) TELEPHONE: (858) 720-2500

65

(B) TELEFAX: (585) 720-2555

66

(C) TELEX: 3760

67

68

(2) INFORMATION FOR SEQ ID NO: 1:

69

70

(i) SEQUENCE CHARACTERISTICS:

71

72

(A) LENGTH: 1785 base pairs

73

(B) TYPE: nucleic acid

74

(C) STRANDEDNESS: single

75

(D) TOPOLOGY: linear

76

77

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

78

79

GGTTATGTCT GACTCACTGC ACTGGAGTTT GGCAAAAGCA TCTCAGAAGT GGTGTGCTT 60

80

TTTTGAATGA AATGATCAAT GGAGTGCTCC AGTTGTATGC TGGCCTCTGG ATACTAACTA 120

81

GACCTGCCTG ACTCCAGGAA CTAAGGCTCA GTATCTGCAG AAGCTTTTTG CCCATCTCAT 180

82

TCCGGCTATG GGGACAACAT GTCTTCACCC AGGAAGGTTA GAGGAAAAAC TGGAAGAGAT 240

83

AATGATGAAG AGGAGGGTAA TTCAGGTAAC CTGAATCTCC GCAACTCTTT GCCTTCATCG 300

84

AGTCAGAAAA TGACGCCTAC GAAGCCGATT TTTGGGAATA AAATGAATTC AGAGAATGTA 360

85

AAACCCTCCC ATCACCTGTC ATTCTCAGAT AAGTATGAGC TTGTTTACCC AGAGCCTTTG 420

86

GAAAGTGACA CTGATGAGAC TGTGTGGGAT GTCAGTGACC GGTCTCTCAG AAACAGGTGG 480

87

AACAGTATGG ATTCAAGAGAC TGCAGGGCCG TCAAAGACTG TCTCCCAGT GCTTCTGGT 540

88

AGTAGTAGGC TCTCAAAGGA CACTGAAACA TCTGTCTCTG AAAAGGAGCT AACTCAGTTG 600

89

GCTCAGATTC GACCATTAAT ATTCAACAGT TCTGCACGGT CTGCTATGCG GGATTGTTT 660

90

AACACGCTTC AGAAAAAAGA AGAACTTGAT ATCATCCGTG AGTTTTTGGA GTTAGAACAA 720

91

ATGACTCTGC CTGATGACTT CAATTCTGGG AATACACTAC AGAACAGAGA TAAGAACAGA 780

92

TACCGAGATA TTCTTCCATA TGATTCAACA CGTGTTCTCTC TTGGAAAAAA CAAGGACTAC 840

93

ATCAACGCTA GTTATATTAG AATAGTAAAT CATGAAGAAG AGTATTTT 900

94

CAAGGACCAT TGCCAGAAAC TATAGAAGAC TTTTGGCAAA TGGTCTTGGA AAATAATTGT 960

95

AATGTTATTG CTATGATAAC CAGAGAGATA GAATGTGGAG TTATCAAGTG TTACAGTTAC 1020

96

TGGCCATTT CTCTGAAGGA GCCTTTGGAA TTCGAACACT TTAGTGTCTT TCTGGAGACC 1080

97

TTTCATGTAA CTCAATATTT CACCGTTCGA GTATTTTCTG TTGTGAAGAA GTCCACAGGA 1140

98

AAGAGCCAAT GTGTAAAACA CTGTCAGTTC ACCAAGTGCG CAGACCATGG CACTCCTGCC 1200

99

TCAGCAGATT TTTTCATAAA ATATGTCCGT TATGTGAGGA AGAGCCACAT TACAGGACCC 1260

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/095,478ADATE: 06/08/2000
TIME: 07:09:27

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101	GTGTTCTCTG	CCATCGAGAA	GAAGTACTCT	TTTGACATTA	TGAACATAGT	GACCCAGATG	1380
102	AGAAAAGCAGC	GCTGTGGCAT	GATTCAAACC	AAGGAGCAGT	ACCAGTTTTG	TTATGAAATT	1440
103	GTGCTTGAAG	TTCTTCAGAA	CCTTCTGGCT	TTGTATTAAG	AGAGACTTCT	GCGCCTGTCC	1500
104	CTCGAGGTTA	CCGAGCAGCT	TGGAGCCTGA	GCCGTGCTGA	AGCGTCTGCG	GGCCGTGCAG	1560
105	TCTGCCTTCT	GATTTTTTCTC	TCTGAAAGTC	CCTGAAGGTA	GCACTACTGG	GCACAGAGTG	1620
106	AACTGTTTCC	ACTTGATCTT	TCTGAACAAG	AGCAAAATAC	CCTCCATGCC	TTCTACGGAA	1680
107	ACGGAAGTTG	CATGAAACAA	CCTCCGCTTG	GCTGTCTGGT	TTGTGGTATT	ACAGAGCTTA	1740
108	ATAAAAGACT	TAGATGTGAA	AAAAAAAAAA	AAAAAAAAAA	AAAAA		1785

109

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111

112 (2) INFORMATION FOR SEQ ID NO:2:

113

114 (i) SEQUENCE CHARACTERISTICS:

115

116 (A) LENGTH: 1896 base pairs

117 (B) TYPE: nucleic acid

118 (C) STRANDEDNESS: single

119 (D) TOPOLOGY: linear

120

121 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

122

123 GGTATATGCTCT GACTCACTGC ACTGGAGTTT GGCAAAAGCA TCTCAGAAGT GGTGTGTGCTT 60

124

125 TTTTGAATGA AATGATCAAT GGAGTGCTCC AGTTGTATGC TGGCCTCTGG ATACTAACTA 120

126 GACCTGCCTG ACTCCAGGAA CTAAGGCTCA GTATCTGCAG AAGCTTTTTG CCCATCTCAT 180

127 TCCGGCTATG GGGACAACAT GTCTTCACCC AGGAAGGTTA GAGGAAAAAC TGGGAAGAGAT 240

128 AATGATGAAG AGGAGGGTAA TTCAGGTAAC CTGAATCTCC GCAACTCTTT GCCTTCATCG 300

129 AGTCAGAAAA TGACGCCTAC GAAGCCGGTA CAAAATAAAA ATCTCATGAA GTATGAAGAA 360

130 CACTTAGATA TATTGATGGT GTTTTTATTG ATAAAAACCA TATGGTATAA TGTCTTCAAA 420

131 TTATGGAAAG GCAAGCTTAT TTTTGGGAAT AAAATGAATT CAGAGAATGT AAAACCTTCC 480

132 CATCACCTGT CATTCTCAGA TAAGTATGAG CTTGTTTACC CAGAGCCTTT GGAAAGTGAC 540

133 ACTGATGAGA CTGTGTGGGA TGTCAGTGAC CGGTCTCTCA GAAACAGGTG GAACAGTATG 600

134 GATTCAAGAGA CTGCAGGGCC GTCAAAGACT GTCTCCCAG TGCTTTCTGG TAGTAGTAGG 660

135 CTCTCAAAGG AACTGAAAC ATCTGTCTCT GAAAAGGAGC TAACTCAGTT GGCTCAGATT 720

136 CGACCATTA TATTCAACAG TTCTGCACGG TCTGCTATGC GGGATTGTTT GAACACGCTT 780

137 CAGAAAAAAG AAGAACTTGA TATCATCCGT GAGTTTTTGG AGTTAGAACA AATGACTCTG 840

138 CCTGATGACT TCAATTCTGG GAATACACTA CAGAACAGAG ATAAGAACAG ATACCGAGAT 900

139 ATTCTTCCAT ATGATTCAAC ACGTGTTCCT CTTGGAAAAA ACAAGGACTA CATCAACGCT 960

140 AGTTATATTA GAATAGTAAA TCATGAAGAA GAGTATTTTT ATATTGCCAC TCAAGGACCA 1020

141 TTGCCAGAAA CTATAGAAGA CTTTGGCAA ATGGTCTGG AAAATAATTG TAATGTTATT 1080

142 GCTATGATAA CCAGAGAGAT AGAATGTGGA GTTATCAAGT GTTACAGTTA CTGGCCCAT 1140

143 TCTCTGAAGG AGCCTTTGGA ATTCGAACAC TTTAGTGTCT TTCTGGAGAC CTTTCATGTA 1200

144 ACTCAATATT TCACCGTTCG AGTATTTTCTG ATTGTGAAGA AGTCCACAGG AAAGAGCCAA 1260

145 TGTGTAATAA ACTTGCAGTT CACCAAGTGG CCAGACCATG GCACTCCTGC CTCAGCAGAT 1320

146 TTTTTCATAA AATATGTCCG TTATGTGAGG AAGAGCCACA TTACAGGACC CCTCCTTGTT 1380

147 CACTGCAGTG CTGGTGTAGG CCGAACAGGG GTGTTCATAT GTGTGGATGT TGTGTTCTCT 1440

148 GCCATCGAGA AGAACTACTC TTTTGACATT ATGAACATAG TGACCCAGAT GAGAAAGCAG 1500

149 CGCTGTGGCA TGATTCAAAC CAAGGAGCAG TACCAGTTTT GTTATGAAAT TGTGCTTGAA 1560

150 GTTCTTCAGA ACCTTCTGGC TTTGTATTAA GAGAGACTTC TGCGCCTGTC CCTCGAGGTT 1620

151 ACCGAGCAGC TTGGAGCCTG AGCCGTGCTG AAGCGTCTGC GGGCCGTGCA GTCTGCCTTC 1680

152 TGATTTTTTCT CTCTGAAAGT CCCTGAAGGT AGCACTACTG GGCACAGAGT GAACTGTTTC 1740

RECEIVED

JUN 26 2000

TECH CENTER 1620/2900

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/095,478ADATE: 06/08/2000
TIME: 07:09:28

INPUT SET: S35575.raw

153	CACTTGATCT	TTCTGAACAA	GAGCAAAATA	CCCTCCATGC	CTTCTACGGA	AACGGAAGTT	1800
154	GCATGAAACA	ACCTCCGCTT	GGCTGTCTGG	TTTGTGGTAT	TACAGAGCTT	AATAAAAGAC	1860
155	TTAGATGTGA	AAAAAAAAAA	AAAAAAAAAA	AAAAAA			1896

156

157

158

159 (2) INFORMATION FOR SEQ ID NO:3:

160

161 (i) SEQUENCE CHARACTERISTICS:

162

163 (A) LENGTH: 1692 base pairs

164 (B) TYPE: nucleic acid

165 (C) STRANDEDNESS: single

166 (D) TOPOLOGY: linear

167

168 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

169

170	GGTTATGTCT	GA	CTCACTGC	ACTGGAGTTT	GGCAAAAGCA	TCTCAGAAAGT	GGTTGTGCTT	60
171	TTTTGAATGA	AATGATCAAT	GGAGTGCTCC	AGTTGTATGC	TGGCCTCTGG	ATACTAACTA		120
172	GACCTGCCTG	ACTCCAGGAA	CTAAGGCTCA	GTATCTGCAG	AAGCTTTTGT	CCCATCTCAT		180
173	TCCGGCTATG	GGGACAACAT	GTCTTCACCC	AGGAAGGTTA	GAGGAAAAAC	TGGAAGAGAT		240
174	AATGATGAAG	AGGAGGGTAA	TTCAGGTAAC	CTGAATCTCC	GCAACTCTTT	GCCTTCATCG		300
175	AGTCAGAAAA	TGACGCCTAC	GAAGCCGATT	TTTGGGAATA	AAATGAATTC	AGAGAATGTA		360
176	AAACCCCTCCC	ATCACCTGTC	ATTCTCAGAT	AAGTATGAGC	TTGTTTACCC	AGAGCCTTTG		420
177	GAAAGTGACA	CTGATGAGAC	TGTGTGGGAT	GTCAGTGACC	GGTCTCTCAG	AAACAGGTGG		480
178	AACAGTATGG	ATTGAGAGAC	TGCAGGGCCG	TCAAAGACTG	TCTCCCAGT	GCTTTCTGGT		540
179	AGTAGTAGGC	TCTCAAAGGA	CACTGAAACA	TCTGTCTCTG	AAAAGGAGCT	AACTCAGTTG		600
180	GCTCAGATTC	GACCATTAAT	ATTCAACAGT	TCTGCACGGT	CTGCTATGCG	GGATTGTTTG		660
181	AACACGCTTC	AGAAAAAAGA	AGAACTTGAT	ATCATCCGTG	AGTTTTTTGA	GTTAGAACAA		720
182	ATGACTCTGC	CTGATGACTT	CAATTCTGGG	AATACACTAC	AGAACAGAGA	TAAGAACAGA		780
183	TACCGAGATA	TTCTTCCATA	TGATTCAACA	CGTGTTCTCT	TTGGAAAAAA	CAAGGACTAC		840
184	ATCAACGCTA	GTTATATTAG	AATAGTAAAT	CATGAAGAAG	AGTATTTTTA	TATTGCCACT		900
185	CAAGGACCAT	TGCCAGAAAC	TATAGAAGAC	TTTTGGCAAA	TGGTTCTGGA	AAATAATTGT		960
186	AATGTTATTG	CTATGATAAC	CAGAGAGATA	GAATGTGGAG	TTATCAAGTG	TTACAGTTAC		1020

187

188	TGGCCCATTT	CTCTGAAGGA	GCCTTTGGAA	TTCGAACACT	TTAGTGTCTT	TCTGGAGACC	1080
189	TTTCATGTAA	CTCAATATTT	CACCGTTCGA	GTATTTTCAGA	TTGTGAAGAA	GTCCACAGGA	1140
190	AAGAGCCAAT	GTGTAAAAACA	CTTGCAAGTTC	ACCAAGTGGC	CAGACCATGG	CACTCCTGCC	1200
191	TCAGCAGATT	TTTTCATAAA	ATATGTCCGT	TATGTGAGGA	AGAGCCACAT	TACAGGACCC	1260
192	CTCCTTGTTT	ACTGCAGTGC	TGGTGTAGGC	CGAACAGGGG	TGTTTCATATG	TGTGGATGTT	1320
193	GTGTTCTCTG	CCATCGAGAA	GA	ACTACTCT	TTTGACATTA	TGAACATAGT	1380
194	AGAAAGCAGC	GCTGTGGCAT	GATTCAAACC	AAGGTTACCG	AGCAGCTTGG	AGCCTGAGCC	1440
195	GTGCTGAAGC	GTCTGCGGGC	CGTGCACTCT	GCCTTCTGAT	TTTTCTCTCT	GAAAGTCCCT	1500
196	GAAGGTAGCA	CTACTGGGCA	CAGAGTGAAC	TGTTTCCACT	TGATCTTTCT	GAACAAGAGC	1560
197	AAAATACCTT	CCATGCCTTC	TACGGAAACG	GAAGTTGCAT	GAAACAACCT	CCGCTTGGCT	1620
198	GTCTGGTTTG	TGGTATTACA	GAGCTTAATA	AAAGACTTAG	ATGTGAAAAA	AAAAAAAAAA	1680
199	AAAAAAAAAA	AA					1692

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201

202

203 (2) INFORMATION FOR SEQ ID NO:4:

204

205 (i) SEQUENCE CHARACTERISTICS:

RAW SEQUENCE LISTING PATENT APPLICATION US/09/095,478A

DATE: 06/08/2000
TIME: 07:09:29

INPUT SET: S35575.raw

206
207 (A) LENGTH: 320 base pairs
208 (B) TYPE: nucleic acid
209 (C) STRANDEDNESS: single
210 (D) TOPOLOGY: linear
211
212 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
213
214 GAAAAATAATT GTAATGTTAT TGCTATGATA ACCAGAGAGA TAGAAGGTGG AGTTATCAAG 60
215 TGTTGCAGTT ACTGGCCCGT TTCTCTGAAG GAGCCTTTGG AATTCAAACA CTTTCATGTC 120
216 CTTCTGGAGA ACTTTCAGAT AACTCAGTAT TTTGTCATCC GAATATTTCA AATTGTGAAG 180
217 AAGTCCACAG GAAAGAGTCA CTCTGTAAAA CACTTGCAGT TCATCAAATG GCCAGACCAT 240
218 GGCACTCCTG CCTCAGTAGA TTTTTCATC AAATATGTCC GTTATGTGAG GAAGAGCCAC 300
219 ATTACAGGAC CCCTCCTTGT 320

220
221
222
223 (2) INFORMATION FOR SEQ ID NO:5:
224

225 (i) SEQUENCE CHARACTERISTICS:
226

227 (A) LENGTH: 426 amino acids
228 (B) TYPE: amino acid
229 (C) STRANDEDNESS: single
230 (D) TOPOLOGY: linear
231

232 (ii) MOLECULE TYPE: peptide
233

234 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
235

236 Met Ser Ser Pro Arg Lys Val Arg Gly Lys Thr Gly Arg Asp Asn Asp
237 1 5 10 15
238
239 Glu Glu Glu Gly Asn Ser Gly Asn Leu Asn Leu Arg Asn Ser Leu Pro
240 20 25 30
241
242 Ser Ser Ser Gln Lys Met Thr Pro Thr Lys Pro Ile Phe Gly Asn Lys
243 35 40 45
244
245 Met Asn Ser Glu Asn Val Lys Pro Ser His His Leu Ser Phe Ser Asp
246 50 55 60
247
248 Lys Tyr Glu Leu Val Tyr Pro Glu Pro Leu Glu Ser Asp Thr Asp Glu
249 65 70 75 80
250
251 Thr Val Trp Asp Val Ser Asp Arg Ser Leu Arg Asn Arg Trp Asn Ser
252 85 90 95
253
254 Met Asp Ser Glu Thr Ala Gly Pro Ser Lys Thr Val Ser Pro Val Leu
255 100 105 110
256
257 Ser Gly Ser Ser Arg Leu Ser Lys Asp Thr Glu Thr Ser Val Ser Glu
258 115 120 125

INPUT SET: S35575.raw

***** PREVIOUSLY ERRORED SEQUENCES - EDITED *****

829 (2) INFORMATION FOR SEQ ID NO:23:

830

831 (i) SEQUENCE CHARACTERISTICS:

832

833 (A) LENGTH: 29 amino acids

834 (B) TYPE: amino acid

835 (C) STRANDEDNESS: single

836 (D) TOPOLOGY: linear

837

838 (ii) MOLECULE TYPE: peptide

839

840 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

841

842 Trp Asp Val Ser Asp Arg Ser Leu Arg Asn Arg Trp Asn Ser Met Asp

843 1 5 10 15

844

845 Ser Glu Thr Ala Gly Pro Ser Lys Thr Val Ser Pro Val

846 20 25

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PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/09/095,478A

DATE: 06/08/2000
TIME: 07:09:31

INPUT SET: S35575.raw

Line

Error

Original Text

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/095,478ADATE: 06/06/2000
TIME: 00:21:16

INPUT SET: S35575.raw

This Raw Listing contains the General
Information Section and those Sequences
containing ERRORS.

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SEQUENCE LISTING

(1) General Information:

(i) APPLICANT: Gregory Plowman
Bahija Jallal

(ii) TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF
PTP RELATED DISORDERS

(iii) NUMBER OF SEQUENCES: 23

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Brobeck, Phleger & Harrison LLP
(B) STREET: 12390 El Camino Real
(C) CITY: San Diego
(D) STATE: California
(E) COUNTRY: U.S.A.
(F) ZIP: 92130-2081

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
storage
(B) COMPUTER: IBM Compatible
(C) OPERATING SYSTEM: IBM P.C. DOS 5.0
(D) SOFTWARE: FastSEQ for Windows 2.0

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: 09/095,478
(B) FILING DATE: June 10, 1998
(C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: 60/049,756

Does Not Comply
Corrected Diskette Needed

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/095,478ADATE: 06/06/2000
TIME: 00:21:17

INPUT SET: S35575.raw

46 (B) FILING DATE: June 11, 1997
47
48 (A) APPLICATION NUMBER:
49 (B) FILING DATE:
50
51
52 (viii) ATTORNEY/AGENT INFORMATION:
53
54 (A) NAME: Warburg, Richard J.
55 (B) REGISTRATION NUMBER: 32,327
56 (C) REFERENCE/DOCKET NUMBER: 235/054
57
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61
62 (ix) TELECOMMUNICATION INFORMATION:
63
64 (A) TELEPHONE: (858) 720-2500
65 (B) TELEFAX: (585) 720-2555
66 (C) TELEX: 3760
67

ERRORED SEQUENCES FOLLOW:

829 (2) INFORMATION FOR SEQ ID NO:23:
830
831 (i) SEQUENCE CHARACTERISTICS:
832
833 (A) LENGTH: 29 amino acids
834 (B) TYPE: amino acid
835 (C) STRANDEDNESS: single
836 (D) TOPOLOGY: linear
837
838 (ii) MOLECULE TYPE: peptide
839
840 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:
841
842 Trp Asp Val Ser Asp Arg Ser Leu Arg Asn Arg Trp Asn Ser Met Asp
843 1 5 10 15
844
845 Ser Glu Thr Ala Gly Pro Ser Lys Thr Val Ser Pro Val
846 20 25
847 110
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852 SSSD/93680. v01
853

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PAGE: 3

RAW SEQUENCE LISTING
PATENT APPLICATION *US/09/095,478A*

DATE: 06/06/2000
TIME: 00:21:18

INPUT SET: S35575.raw

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PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/09/095,478A

DATE: 06/06/2000
TIME: 00:21:19

INPUT SET: S35575.raw

Line	Error	Original Text
852	Wrong Amino Acid Designator	SSSD/93680. v01
852	Wrong Amino Acid Designator	SSSD/93680. v01